



SEQUENCE LISTING

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<120> RPS2 GENE FAMILY, PRIMERS, PROBES, AND
DETECTION METHODS

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<141> 2003-07-02

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His Ser His Lys Thr Arg Asp Tyr Val Ile Ile Lys Thr Lys Leu Ser
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Tyr Arg Trp Leu Cys Ser Gly Val Val
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Leu Glu Gly His Thr
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Glu His Trp			
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 Gln Arg Ala Asp His Lys Arg Thr Ser Val
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<210> 78

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Glu Asn Thr Gly Glu Asp Ala
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Lys Thr His Met Pro Glu Thr Asp Asn Thr Asp Ala Pro Thr Glu Gly
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Gln Asn Pro Asp Leu Ala
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35 40 45
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gctccagttg ccataaatca cagcccgtc agcagggagg tcccgtcaca cgcggcaccc 240
actcaggcaa agcaaacc aa ccttcaatct gaagctggcg atttagatgc aagaaaaagt 300
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Ser Glu Ala Gly Asp Leu Asp Ala Arg Lys Ser Ser Ala Ser Ser Pro
          35           40           45
Glu Thr Arg Ala Leu Leu Ala Thr Lys Thr Val Leu Gly Arg His Lys
          50           55           60
Ile Glu Val Pro Ala Phe Gly Gly Trp Phe Lys Lys Lys Ser Ser Lys
          65           70           75           80
His Glu Thr Gly Gly Ser Ser Ala Asn Ala Asp Ser Ser Ser Val Ala
          85           90           95
Ser Asp Ser Thr Glu Lys Pro Leu Phe Arg Leu Thr His Val Pro Tyr
          100          105          110
Val Ser Gln Gly Asn Glu Arg Met Gly Cys Trp Tyr Ala Cys Ala Arg
          115          120          125
Met Val Gly His Ser Val Glu Ala Gly Pro Arg Leu Gly Leu Pro Glu
          130          135          140
Leu Tyr Glu Gly Arg Glu Ala Pro Ala Gly Leu Gln Asp Phe Ser Asp
          145          150          155          160
Val Glu Arg Phe Ile His Asn Glu Gly Leu Thr Arg Val Asp Leu Pro
          165          170          175
Asp Asn Glu Arg Phe Thr His Glu Glu Leu Gly Ala Leu Leu Tyr Lys
          180          185          190
His Gly Pro Ile Ile Phe Gly Trp Lys Thr Pro Asn Asp Ser Trp His
          195          200          205
Met Ser Val Leu Thr Gly Val Asp Lys Glu Thr Ser Ser Ile Thr Phe
          210          215          220
His Asp Pro Arg Gln Gly Pro Asp Leu Ala Met Pro Leu Asp Tyr Phe
          225          230          235          240
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Val	Asn	Asp	Asp	Asp	Asp	Ser	Thr	Ser	Glu	Val	Asp	Ala	Ile	Ser	Asp
		35					40					45			
Ser	Thr	Asn	Pro	Ser	Gly	Ser	Phe	Pro	Ser	Val	Glu	Tyr	Glu	Val	Phe
	50					55					60				
Leu	Ser	Phe	Arg	Gly	Pro	Asp	Thr	Arg	Glu	Gln	Phe	Thr	Asp	Phe	Leu
65				70						75					80
Tyr	Gln	Ser	Leu	Arg	Arg	Tyr	Lys	Ile	His	Thr	Phe	Arg	Asp	Asp	Asp
				85					90					95	
Glu	Leu	Leu	Lys	Gly	Lys	Glu	Ile	Gly	Pro	Asn	Leu	Leu	Arg	Ala	Ile
			100					105						110	
Asp	Gln	Ser	Lys	Ile	Tyr	Val	Pro	Ile	Ile	Ser	Ser	Gly	Tyr	Ala	Asp
		115					120						125		
Ser	Lys	Trp	Cys	Leu	Met	Glu	Leu	Ala	Glu	Ile	Val	Arg	Arg	Gln	Glu
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Glu	Asp	Pro	Arg	Arg	Ile	Ile	Leu	Pro	Ile	Phe	Tyr	Met	Val	Asp	Pro
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Ser	Asp	Val	Arg	His	Gln	Thr	Gly	Cys	Tyr	Lys	Lys	Ala	Phe	Arg	Lys
				165					170					175	
His	Ala	Asn	Lys	Phe	Asp	Gly	Gln	Thr	Ile	Gln	Asn	Trp	Lys	Asp	Ala
			180					185						190	
Leu	Lys	Lys	Val	Gly	Asp	Leu	Lys	Gly	Trp	His	Ile	Gly	Lys	Asn	Asp
		195					200							205	
Lys	Gln	Gly	Ala	Ile	Ala	Asp	Lys	Val	Ser	Ala	Asp	Ile	Trp	Ser	His
	210					215					220				
Ile	Ser	Lys	Glu	Asn	Leu	Ile	Leu	Glu	Thr	Asp	Glu	Leu	Val	Gly	Ile
225				230						235					240
Asp	Asp	His	Ile	Thr	Ala	Val	Leu	Glu	Lys	Leu	Ser	Leu	Asp	Ser	Glu
				245					250					255	
Asn	Val	Thr	Met	Val	Gly	Leu	Tyr	Gly	Met	Gly	Gly	Ile	Gly	Lys	Thr
			260					265						270	
Thr	Thr	Ala	Lys	Ala	Val	Tyr	Asn	Lys	Ile	Ser	Ser	Cys	Phe	Asp	Cys
		275					280						285		
Cys	Cys	Phe	Ile	Asp	Asn	Ile	Arg	Glu	Thr	Gln	Glu	Lys	Asp	Gly	Val
	290					295					300				
Val	Val	Leu	Gln	Lys	Lys	Leu	Val	Ser	Glu	Ile	Leu	Arg	Ile	Asp	Ser
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Gly	Ser	Val	Gly	Phe	Asn	Asn	Asp	Ser	Gly	Gly	Arg	Lys	Thr	Ile	Lys
				325					330					335	
Glu	Arg	Val	Ser	Arg	Phe	Lys	Ile	Leu	Val	Val	Leu	Asp	Asp	Val	Asp
			340					345					350		
Glu	Lys	Phe	Lys	Phe	Glu	Asp	Met	Leu	Gly	Ser	Pro	Lys	Asp	Phe	Ile
		355					360					365			
Ser	Gln	Ser	Arg	Phe	Ile	Ile	Thr	Ser	Arg	Ser	Met	Arg	Val	Leu	Gly
	370					375					380				
Thr	Leu	Asn	Glu	Asn	Gln	Cys	Lys	Leu	Tyr	Glu	Val	Gly	Ser	Met	Ser
385					390					395					400
Lys	Pro	Arg	Ser	Leu	Glu	Leu	Phe	Ser	Lys	His	Ala	Phe	Lys	Lys	Asn
				405					410					415	
Thr	Pro	Pro	Ser	Ser	Tyr	Tyr	Glu	Thr	Leu	Ala	Asn	Asp	Val	Val	Asp
			420					425					430		
Thr	Thr	Ala	Gly	Leu	Pro	Leu	Thr	Leu	Lys	Val	Ile	Gly	Ser	Leu	Leu

[illegible]

Pro	Thr	Trp	Leu	Pro	Gly	Ile	Glu	Asn	Leu	Glu	Asn	Leu	Thr	Ser	Leu		
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		915					920					925					
Gln	Gly	Leu	Arg	Ser	Leu	Glu	Ile	Leu	Arg	Ile	Arg	Lys	Val	Asn	Gly		
	930					935					940						
Leu	Ala	Arg	Ile	Lys	Gly	Leu	Lys	Asp	Leu	Leu	Cys	Ser	Ser	Thr	Cys		
945					950					955					960		
Lys	Leu	Arg	Lys	Phe	Tyr	Ile	Thr	Glu	Cys	Pro	Asp	Leu	Ile	Glu	Leu		
			965					970						975			
Leu	Pro	Cys	Glu	Leu	Gly	Val	Gln	Thr	Val	Val	Val	Pro	Ser	Met	Ala		
		980					985						990				
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	995						1000					1005					
Arg	Ser	Leu	Pro	Lys	Phe	Pro	Met	Leu	Lys	Lys	Leu	Asp	Leu	Ala	Val		
	1010					1015					1020						
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Glu	Leu	Val	Ser	Leu	Glu	Leu	Glu	Leu	Asp	Asp	Thr	Ser	Ser	Gly	Ile		
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Val	Lys	Val	Pro	Ser	Leu	Arg	Glu	Ile	Glu	Gly	Leu	Glu	Glu	Leu	Lys		
	1075					1080						1085					
Ser	Leu	Gln	Asp	Leu	Tyr	Leu	Glu	Gly	Cys	Thr	Ser	Leu	Gly	Arg	Leu		
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Thr	Glu	Leu	Val	Gln	Thr	Val	Val	Ala	Val	Pro	Ser	Leu	Arg	Gly	Leu		
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Thr	Ile	Arg	Asp	Cys	Pro	Arg	Leu	Glu	Val	Gly	Pro	Met	Ile	Gln	Ser		
		1140						1145					1150				
Leu	Pro	Lys	Phe	Pro	Met	Leu	Asn	Glu	Leu	Thr	Leu	Ser	Met	Val	Asn		
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Ile	Thr	Lys	Glu	Asp	Glu	Leu	Glu	Val	Leu	Gly	Ser	Leu	Glu	Glu	Leu		
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Ile	Ser	Phe	Leu	Ser	Lys	Leu	Gln	Lys	Leu	Thr	Thr	Leu	Ile	Val	Glu		
			1205					1210						1215			
Val	Pro	Ser	Leu	Arg	Glu	Ile	Glu	Gly	Leu	Ala	Glu	Leu	Lys	Ser	Leu		
		1220						1225				1230					
Arg	Ile	Leu	Tyr	Leu	Glu	Gly	Cys	Thr	Ser	Leu	Glu	Arg	Leu	Trp	Pro		
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		35					40					45					
Leu	Glu	Tyr	Gly	Ala	Thr	Ile	Pro	Gly	Glu	Leu	Cys	Lys	Ala	Ile	Glu		
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65					70					75					80		
Arg	Trp	Cys	Leu	Asn	Glu	Leu	Val	Lys	Ile	Met	Glu	Cys	Lys	Thr	Arg		
				85					90					95			
Phe	Lys	Gln	Thr	Val	Ile	Pro	Ile	Phe	Tyr	Asp	Val	Asp	Pro	Ser	His		
			100					105					110				
Val	Arg	Asn	Gln	Lys	Glu	Ser	Phe	Ala	Lys	Ala	Phe	Glu	Glu	His	Glu		
		115					120					125					
Thr	Lys	Tyr	Lys	Asp	Asp	Val	Glu	Gly	Ile	Gln	Arg	Trp	Arg	Ile	Ala		
	130					135					140						
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 35 40 45
 Arg Asp Asp Leu Thr Leu Arg Ile Gln Gln Asp Gly Leu Glu Gly Arg
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 Ser Cys Ser Asn Arg Ala Arg Glu Trp Leu Ser Ala Val Gln Val Thr
 65 70 75 80
 Glu Thr Lys Thr Ala Leu Leu Leu Val Arg Phe Arg Arg Arg Glu Gln
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 Arg Thr Arg Met Arg Arg Arg Tyr Leu Ser Cys Phe Gly Cys Ala Asp
 100 105 110
 Tyr Lys Leu Cys Lys Lys Val Ser Ala Ile Leu Lys Ser Ile Gly Glu
 115 120 125
 Leu Arg Glu Arg Ser Glu Ala Ile Lys Thr Asp Gly Gly Ser Ile Gln
 130 135 140
 Val Thr Cys Arg Glu Ile Pro Ile Lys Ser Val Val Gly Asn Thr Thr
 145 150 155 160
 Met Met Glu Gln Val Leu Glu Phe Leu Ser Glu Glu Glu Glu Arg Gly
 165 170 175
 Ile Ile Gly Val Tyr Gly Pro Gly Gly Val Gly Lys Thr Thr Leu Met
 180 185 190
 Gln Ser Ile Asn Asn Glu Leu Ile Thr Lys Gly His Gln Tyr Asp Val
 195 200 205
 Leu Ile Trp Val Gln Met Ser Arg Glu Phe Gly Glu Cys Thr Ile Gln
 210 215 220
 Gln Ala Val Gly Ala Arg Leu Gly Leu Ser Trp Asp Glu Lys Glu Thr
 225 230 235 240
 Gly Glu Asn Arg Ala Leu Lys Ile Tyr Arg Ala Leu Arg Gln Lys Arg
 245 250 255
 Phe Leu Leu Leu Leu Asp Asp Val Trp Glu Glu Ile Asp Leu Glu Lys
 260 265 270
 Thr Gly Val Pro Arg Pro Asp Arg Glu Asn Lys Cys Lys Val Met Phe
 275 280 285
 Thr Thr Arg Ser Ile Ala Leu Cys Asn Asn Met Gly Ala Glu Tyr Lys
 290 295 300
 Leu Arg Val Glu Phe Leu Glu Lys Lys His Ala Trp Glu Leu Phe Cys
 305 310 315 320
 Ser Lys Val Trp Arg Lys Asp Leu Leu Glu Ser Ser Ser Ile Arg Arg
 325 330 335
 Leu Ala Glu Ile Ile Val Ser Lys Cys Gly Gly Leu Pro Leu Ala Leu
 340 345 350
 Ile Thr Leu Gly Gly Ala Met Ala His Arg Glu Thr Glu Glu Glu Trp
 355 360 365

Ile	His	Ala	Ser	Glu	Val	Leu	Thr	Arg	Phe	Pro	Ala	Glu	Met	Lys	Gly	370	375	380
Met	Asn	Tyr	Val	Phe	Ala	Leu	Leu	Lys	Phe	Ser	Tyr	Asp	Asn	Leu	Glu	385	390	395
Ser	Asp	Leu	Leu	Arg	Ser	Cys	Phe	Leu	Tyr	Cys	Ala	Leu	Phe	Pro	Glu	405	410	415
Glu	His	Ser	Ile	Glu	Ile	Glu	Gln	Leu	Val	Glu	Tyr	Trp	Val	Gly	Glu	420	425	430
Gly	Phe	Leu	Thr	Ser	Ser	His	Gly	Val	Asn	Thr	Ile	Tyr	Lys	Gly	Tyr	435	440	445
Phe	Leu	Ile	Gly	Asp	Leu	Lys	Ala	Ala	Cys	Leu	Leu	Glu	Thr	Gly	Asp	450	455	460
Glu	Lys	Thr	Gln	Val	Lys	Met	His	Asn	Val	Val	Arg	Ser	Phe	Ala	Leu	465	470	475
Trp	Met	Ala	Ser	Glu	Gln	Gly	Thr	Tyr	Lys	Glu	Leu	Ile	Leu	Val	Glu	485	490	495
Pro	Ser	Met	Gly	His	Thr	Glu	Ala	Pro	Lys	Ala	Glu	Asn	Trp	Arg	Gln	500	505	510
Ala	Leu	Val	Ile	Ser	Leu	Leu	Asp	Asn	Arg	Ile	Gln	Thr	Leu	Pro	Glu	515	520	525
Lys	Leu	Ile	Cys	Pro	Lys	Leu	Thr	Thr	Leu	Met	Leu	Gln	Gln	Asn	Ser	530	535	540
Ser	Leu	Lys	Lys	Ile	Pro	Thr	Gly	Phe	Phe	Met	His	Met	Pro	Val	Leu	545	550	555
Arg	Val	Leu	Asp	Leu	Ser	Phe	Thr	Ser	Ile	Thr	Glu	Ile	Pro	Leu	Ser	565	570	575
Ile	Lys	Tyr	Leu	Val	Glu	Leu	Tyr	His	Leu	Ser	Met	Ser	Gly	Thr	Lys	580	585	590
Ile	Ser	Val	Leu	Pro	Gln	Glu	Leu	Gly	Asn	Leu	Arg	Lys	Leu	Lys	His	595	600	605
Leu	Asp	Leu	Gln	Arg	Thr	Gln	Phe	Leu	Gln	Thr	Ile	Pro	Arg	Asp	Ala	610	615	620
Ile	Cys	Trp	Leu	Ser	Lys	Leu	Glu	Val	Leu	Asn	Leu	Tyr	Tyr	Ser	Tyr	625	630	635
Ala	Gly	Trp	Glu	Leu	Gln	Ser	Phe	Gly	Glu	Asp	Glu	Ala	Glu	Glu	Leu	645	650	655
Gly	Phe	Ala	Asp	Leu	Glu	Tyr	Leu	Glu	Asn	Leu	Thr	Thr	Leu	Gly	Ile	660	665	670
Thr	Val	Leu	Ser	Leu	Glu	Thr	Leu	Lys	Thr	Leu	Phe	Glu	Phe	Gly	Ala	675	680	685
Leu	His	Lys	His	Ile	Gln	His	Leu	His	Val	Glu	Glu	Cys	Asn	Glu	Leu	690	695	700
Leu	Tyr	Phe	Asn	Leu	Pro	Ser	Leu	Thr	Asn	His	Gly	Arg	Asn	Leu	Arg	705	710	715
Arg	Leu	Ser	Ile	Lys	Ser	Cys	His	Asp	Leu	Glu	Tyr	Leu	Val	Thr	Pro	725	730	735
Ala	Asp	Phe	Glu	Asn	Asp	Trp	Leu	Pro	Ser	Leu	Glu	Val	Leu	Thr	Leu	740	745	750
His	Ser	Leu	His	Asn	Leu	Thr	Arg	Val	Trp	Gly	Asn	Ser	Val	Ser	Gln	755	760	765
Asp	Cys	Leu	Arg	Asn	Ile	Arg	Cys	Ile	Asn	Ile	Ser	His	Cys	Asn	Lys	770	775	780
Leu	Lys	Asn	Val	Ser	Trp	Val	Gln	Lys	Leu	Pro	Lys	Leu	Glu	Val	Ile	785	790	795
Glu	Leu	Phe	Asp	Cys	Arg	Glu	Ile	Glu	Glu	Leu	Ile	Ser	Glu	His	Glu	805	810	815
Ser	Pro	Ser	Val	Glu	Asp	Pro	Thr	Leu	Phe	Pro	Ser	Leu	Lys	Thr	Leu			

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<400> 144
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<211> 26

<212> DNA

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<222> 3, 9, 12, 13, 15, 18, 24

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13

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<223> n = A,T,C or G

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<210> 177
<211> 20
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<400> 177
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<210> 178
<211> 23
<212> DNA
<213> Arabidopsis thaliana

<220>
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<400> 178
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<210> 179
<211> 17
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<211> 21
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<220>
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<220>
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15

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21

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17

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<400> 188
Leu Lys Phe Ser Tyr Asp Asn Leu Glu Ser Asp Leu Leu
1 5 10

<210> 189
<211> 16
<212> PRT
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<400> 189
Gly Val Tyr Gly Pro Gly Gly Val Gly Lys Thr Thr Leu Met Gln Ser
1 5 10 15

<210> 190
<211> 14
<212> PRT
<213> Arabidopsis thaliana

<400> 190
Gly Gly Leu Pro Leu Ala Leu Ile Thr Leu Gly Gly Ala Met
1 5 10

<210> 191
<211> 11
<212> PRT

<213> Arabidopsis thaliana

<220>

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<221> VARIANT

<222> 3

<223> Xaa = Gly or Pro

<221> VARIANT

<222> 5

<223> Xaa = Ile, Leu, or Val

<221> VARIANT

<222> 10

<223> Xaa = Ile, Leu, or Thr

<221> VARIANT

<222> 11

<223> Xaa = Ala or Met

<400> 191

Gly Xaa Xaa Gly Xaa Gly Lys Thr Thr Xaa Xaa
1 5 10

<210> 192

<211> 11

<212> PRT

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<221> VARIANT

<222> 2

<223> Xaa = Arg or Lys

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<222> 3

<223> Xaa = Ile, Val, or Phe

<221> VARIANT

<222> 5

<223> Xaa = Ile, Leu, or Val

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<222> 6

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<222> 7

<223> Xaa = Ile or Val

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<223> Xaa = Ile , Leu, or Val

<221> VARIANT
<222> (11)...(11)
<223> Xaa = Asp or Trp

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1 5 10

<210> 193
<211> 8
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<223> Xaa = Arg or Lys

<221> VARIANT
<222> 3
<223> Xaa = Phe, Ile, or Val

<221> VARIANT
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<223> Xaa = Ile or Met

<221> VARIANT
<222> 5
<223> Xaa = Ile, Leu, or Phe

<221> VARIANT
<222> 7
<223> Xaa = Ser, Cys, or Thr

<400> 193
Xaa Xaa Xaa Xaa Xaa Thr Xaa Arg
1 5

<210> 194
<211> 8
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<213> Arabidopsis thaliana

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<223> Xaa = Thr, Ala, or Ser

<221> VARIANT

<222> 6

<223> Xaa = Leu or Val

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<222> 7

<223> Xaa = Ile, Val, or Lys

<221> VARIANT

<222> 8

<223> Xaa = Val or Thr

<400> 194

Gly Leu Pro Leu Xaa Xaa Xaa Xaa

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5

<210> 195

<211> 7

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<223> Xaa = Ile or Phe

<221> VARIANT

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<223> Xaa = Asp or Lys

<221> VARIANT

<222> 6

<223> Xaa = Ala, Gly, or Asn

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5

<210> 196

<211> 4

<212> PRT

<213> Arabidopsis thaliana

<400> 196

Asn Ser His Arg

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<210> 197

<400> 197
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<210> 198
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<400> 198
Thr Gly Asp Leu
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<210> 199
<211> 4
<212> PRT
<213> Arabidopsis thaliana

<400> 199
His Gly Thr Tyr
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<210> 200
<211> 11
<212> PRT
<213> Arabidopsis thaliana

<400> 200
Arg Met Ser His Gly Phe Arg Asn Ser Gln Ser
1 5 10

<210> 201
<211> 27
<212> PRT
<213> Arabidopsis thaliana

<400> 201
Gly Glu Met Val Glu Ser Thr Gly Lys Arg Ser Thr Lys Arg Arg Ala
1 5 10 15
Leu Leu Phe Thr Ala Leu Cys Ser Lys Leu Ile
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<210> 202
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<212> PRT
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<220>
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<211> 6
<212> PRT
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<220>
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1 5

<210> 204
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<223> Xaa = Gln or Leu

<221> VARIANT
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<223> Xaa = Leu or Ile

<221> VARIANT
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<400> 204
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1 5

<210> 205
<211> 6
<212> PRT
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<400> 205
Ser Lys Leu Lys Ser Leu
1 5

<210> 206
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<213> Arabidopsis thaliana

<220>
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<223> Xaa = Arg or His

<221> VARIANT
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<223> Xaa = Ile or Tyr

<400> 206
Gly Leu Xaa Ser Leu Glu Xaa Leu
1 5

<210> 207
<211> 6
<212> PRT
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<400> 207
Ser Lys Leu Lys Ser Leu
1 5

<210> 208
<211> 7
<212> PRT
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<400> 208
Lys Phe Ser Tyr Asp Asn Leu
1 5

<210> 209
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<220>
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<223> Xaa = Any Amino Acid

<221> VARIANT
<222> 4, 15, 20, 23
<223> Xaa = Leu, Ile, or Val

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1 5 10 15
Xaa Xaa Xaa Xaa Xaa Xaa Xaa
20

<210> 210
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<212> PRT
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 <220>
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 <222> 2, 3, 5, 6, 8, 9, 11, 12, 14, 16, 17, 19, 21, 22
 <223> Xaa = Any Amino Acid

 <221> VARIANT
 <222> 4, 20, 23
 <223> Xaa = Leu, Ile, or Val

 <400> 210
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 1 5 10 15
 Xaa Asn Xaa Xaa Xaa Xaa Xaa
 20

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 <220>
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 <222> 2, 3, 5, 6, 8, 9, 11
 <223> Xaa = Any Amino Acid

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 <222> 1
 <223> Xaa = Ile, Leu, or Val

 <221> VARIANT
 <222> 10
 <223> Xaa = Ile or Leu

 <400> 211
 Xaa Xaa Xaa Leu Xaa Xaa Leu Xaa Xaa Xaa Xaa Leu
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 <210> 212
 <211> 7
 <212> PRT
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 <220>
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 <223> Xaa = Ile or Arg

 <221> VARIANT
 <222> 2, 5-7
 <223> Xaa = Any Amino Acid

 <400> 212

Xaa Xaa Asp Leu Xaa Xaa Xaa
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<210> 213
<211> 8
<212> PRT
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<400> 213
Gly Pro Gly Gly Val Gly Lys Thr
1 5

<210> 214
<211> 16
<212> PRT
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<400> 214
Thr Tyr Gly Ala Tyr Gly Ala Tyr Arg Thr Asx Tyr Arg Asx Arg Ala
1 5 10 15

<210> 215
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<212> DNA
<213> Arabidopsis thaliana

<400> 215
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gccatcggtg acttgaaggc catacgtgat gacctgactt tacggatcca acaagacggt 180
ctagagggac gaagctgctc aaatcgtgcc agagagtggc ttagtgcggt gcaagtaacg 240
gagactaaaa cagccctact tttagttagg tttaggcgctc gggaacagag gacgcgaatg 300
aggaggagat acctcagttg tttcggttgt gccgactaca aactgtgcaa gaaggtttct 360
gccatattga agagcattgg tgagctgaga gaacgctctg aagctatcaa aacagatggc 420
gggtcaattc aagtaacttg tagagagata cccatcaagt ccggtgtcgg aaataccacg 480
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acaaaaggac atcagtatga tgtactgatt tgggttcaaa tgtccagaga attcggcgag 660
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catactgaag ctcttaaagc agaaaactgg cgacaagcgt tgggtgatctc attgttagat 1560
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ttggaatact	tggaaaacct	aaccacactc	ggtatcactg	ttctctcatt	ggagacccta	2040
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tgcaatgaac	tcctctactt	caatctccca	tcactcacta	accatggcag	gaacctgaga	2160
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gaagatccaa	cattgttccc	aagcctgaag	accttgagaa	ctagggatct	gccagaacta	2520
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tgccccagag	ttaagaaact	gccgtttcag	gagaggagga	cccagatgaa	cttgccaaca	2640
gtttattgtg	aggagaaatg	gtggaaagca	ctggaaaaag	atcaaccaa	cgaagagctt	2700
tgttatttac	cgcgctttgt	tccaaattga				2730

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 <213> Arabidopsis thaliana

<400> 216	
gtggagttgt atg	13

<210> 217
 <211> 6
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 <213> Arabidopsis thaliana

<400> 217
 Gly Leu Pro Leu Ala Leu
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